IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Applicants:	Baker et al.	Docket No:	39780-2830P1C7
Serial No:	10/006,130	Group Art Unit:	1647
Filed:	December 6, 2001	Examiner:	Rachel K. Hunnicutt

For: **SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC ACIDS ENCODING THE SAME**

Commissioner for Patents
P.O. Box 1450
Alexandria, Virginia 22313-1450

DECLARATION OF NAPOLEONE FERRARA, Ph.D., AUDREY GODDARD, Ph.D., PAUL J. GODOWSKI, Ph.D., AUSTIN GURNEY, Ph.D., JAMES PAN, Ph.D., COLIN K. WATANABE and WILLIAM I. WOOD, Ph.D. UNDER 37 CFR 1.131

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10. The DNA-64883 sequence shown in the GSeqEdit report is identical to that of SEQ ID NO: 129 disclosed in the above-identified application.

11. The beginning of the cDNA sequence corresponding to SEQ ID NO: 129 in the above-identified application is shown on page 1 of the GSeqEdit database report, and the location of the first nucleotide is marked with "^insert starts here" and an arrow. The location of the last nucleotide corresponding to SEQ ID NO: 129 is shown on page 11 and is marked with an arrow.

12. The amino acid sequence shown in the GSeqEdit report is identical to that of SEQ ID NO: 130 disclosed in the above-identified application.

13. The first 26 amino acid residues of the PRO1244 polypeptide (SEQ ID NO:130) encoded by the cDNA (DNA-64883) are also shown on page 1 of the GSeqEdit report and the remaining 309 residues appear on pages 2-6 of the report.

14. All activities listed under paragraphs 4-13 were completed prior to August 14, 1998. (See Exhibit A).

15. We hereby declare that all statements made herein of our own knowledge are true and that all statements made on information or belief are believed to be true, and further that these statements were made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United States Code and that such willful statements may jeopardize the validity of the application or any patent issued thereon.

Napoleone Ferrara, Ph.D.	Date
Audrey Goddard, Ph.D.	Date
Paul J. Godowski, Ph.D.	Date
James Pan, Ph.D.	Date
Austin Gurney, Ph.D.	Date
Colin K. Watanabe	Date
William I. Wood, Ph.D.	10/5/04 Date

SV 2060299 v2
10/1/04 2:02 PM (39780.2830)

>[redacted]
>DNA64883 [Full]
>510 Sites [All Sites]
> [redacted] DNA64883 wiw GSeqEdit
> [redacted] DNA64883 zemin GSeqEdit
> [redacted] DNA64883 goddarda GSeqEdit
> [redacted] DNA64883 sheldens GSeqEdit
>HBN64883.seq, sequenced at ABI/ACGT by Peter Ma and Ellson Chen
>human ortholog of implantation-associated protein - Rattus

```
                                                                   nlaIII
              mnlI                                                 mslI
             taqI                                                 styI
            xhoI              thaI                                ncoI
            tliI              fnuDII/mvnI                         dsaI         tseI
            smlI             hinPI   aciI                         btgI/bstDSI  fnu4HI/bsoFI
            paeR7I  mwoI   tseI bstUI[M.hhaI-]                    bstXI        bbvI                 dde
       tsp509I[M.ecoRI-]   fnu4HI/bsoFI                           bsaJI       hinPI            taiI   bsp
      ecoRI     avaI[M.taqI-]  bbvI bsh1236I                   tsp45I         hhaI/cfoI        maeII/hpyC
      apoI   mwoI bseRI   nlaIII  hhaI/cfoI                 bsmAI maeIII     haeII          hpy99I   mnlI
 1 CGGAATTCGG CTCGAGGAGC GAACATGGCA GCGCGTTGGC GGTTTTGGTG TGTCTCTGTG ACCATGGTGG TGGCGCTGCT CATCGTTTGC GACGTTCCCT
   GCCTTAAGCC GAGCTCCTCG CTTGTACCGT CGCGCAACCG CCAAAACCAC ACAGAGACAC TGGTACCACC ACCGCGACGA GTAGCAAACG CTGCAAGGGA
 1                         M  A  A  R  W  R  F  W  C  V  S  V  T  M  V  V  A  L  L  I  V  C  D  V  P  S
                ^insert starts here
                           ^MET
```

```
     mnlI                                                   aluI
    alwNI[dcm-]                                            pvuII
    alw26I/bsmAI          bsaXI      hpy188I               mspA1I/nspBII                                      bsmAI
101 CAGCCTCTGC CCAAAGAAAG AAGGAGATGG TGTTATCTGA AAAGGTTAGT CAGCTGATGG AATGGACTAA CAAAAGACCT GTAATAAGAA TGAATGGAGA
    GTCGGAGACG GGTTTCTTTC TTCCTCTACC ACAATAGACT TTTCCAATCA GTCGACTACC TTACCTGATT GTTTTCTGGA CATTATTCTT ACTTACCTCT
27   A  S  A   Q  R  K   K  E  M  V   L  S  E   K  V  S   Q  L  M  E   W  T  N   K  R  P   V  I  R  M   N  G  D

                                                                  tspRI                 bst4CI/hpyCH4III   cac8I
                                                                  btsI                 ahdI/eam1105I    cac8I
        hpy99I                        tsp509I             nlaIII                    hpyCH4V tspRI        hpyCH4V al
201 CAAGTTCCGT CGCCTTGTGA AAGCCCCACC GAGAAATTAC TCCGTTATCG TCATGTTCAC TGCTCTCCAA CTGCATAGAC AGTGTGTCGT TTGCAAGCAA
    GTTCAAGGCA GCGGAACACT TTCGGGGTGG CTCTTTAATG AGGCAATAGC AGTACAAGTG ACGAGAGGTT GACGTATCTG TCACACAGCA AACGTTCGTT
60   K  F  R   R  L  V  K   A  P  P   R  N  Y   S  V  I  V   M  F  T   A  L  Q   L  H  R  Q   C  V  V   C  K  Q
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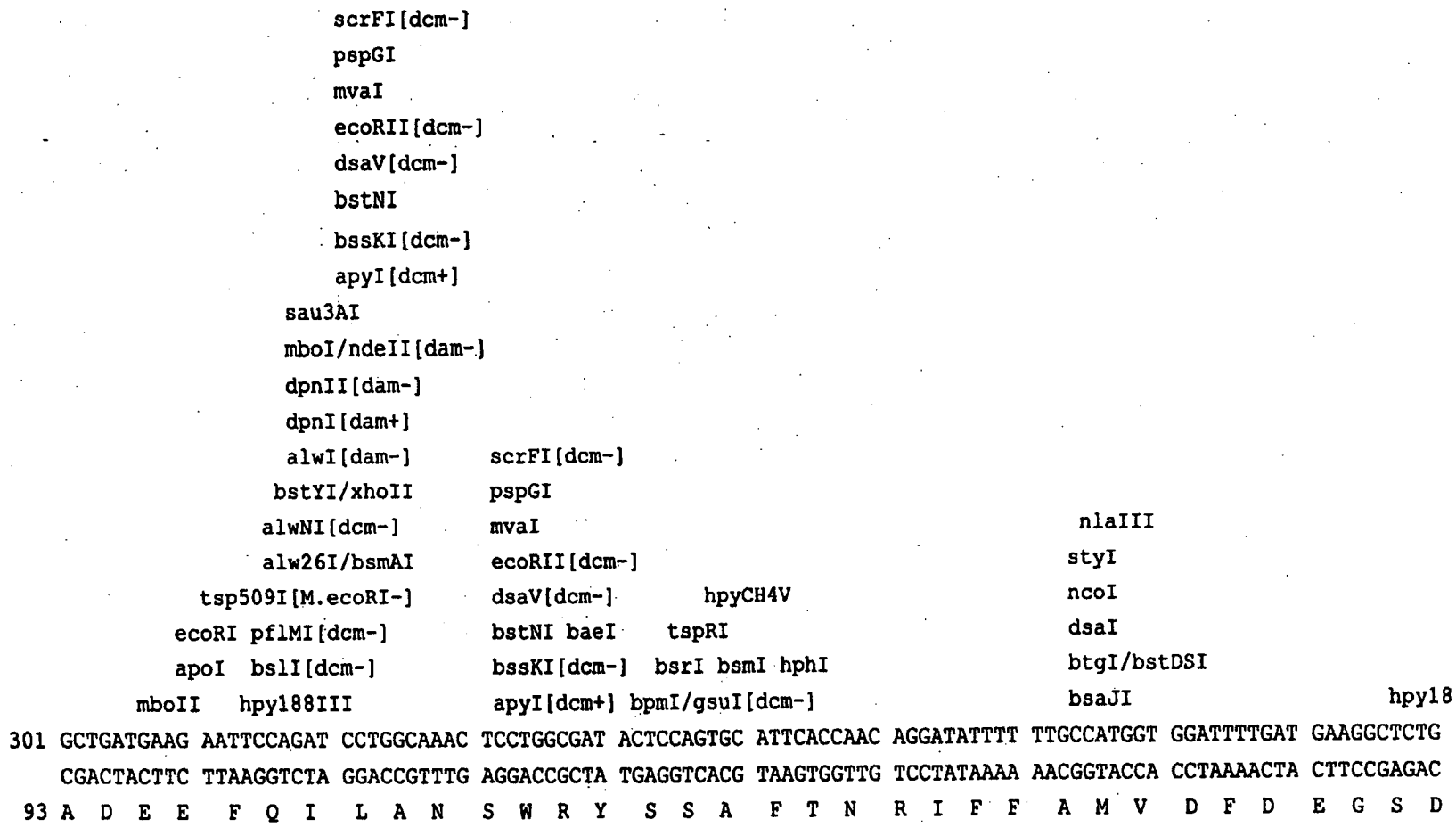
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pspGI
mvaI
ecoRII[dcm-]
dsaV[dcm-]
bstNI
bssKI[dcm-]
apyI[dcm+]
sau3AI
mboI/ndeII[dam-]
dpnII[dam-]
dpnI[dam+]
alwI[dam-]
bstYI/xhoII
alwNI[dcm-]
alw26I/bsmAI
tsp509I[M.ecoRI-]
ecoRI pflMI[dcm-]
apoI bslI[dcm-]
mboII hpy188III
scrFI[dcm-]
pspGI
mvaI
ecoRII[dcm-]
dsaV[dcm-]
bstNI baeI
bssKI[dcm-]
apyI[dcm+]
hpyCH4V
tspRI
bsrI bsmI hphI
bpmI/gsuI[dcm-]
nlaIII
styI
ncoI
dsaI
btgI/bstDSI
bsaJI
hpy18
301 GCTGATGAAG AATTCCAGAT CCTGGCAAAC TCCTGGCGAT ACTCCAGTGC ATTCACCAAC AGGATATTTT TTGCCATGGT GGATTTTGAT GAAGGCTCTG
CGACTACTTC TTAAGGTCTA GGACCGTTTG AGGACCGCTA TGAGGTCACG TAAGTGGTTG TCCTATAAAA AACGGTACCA CCTAAAACTA CTTCCGAGAC
93 A D E E F Q I L A N S W R Y S S A F T N R I F F A M V D F D E G S D

```
                         tsp509I[M.ecoRI-]
                         ecoRI                                   hpyCH4V
              sfaNI      apoI                                   ecoNI
           hpy188I    nlaIII   aluI                             bslI       bslI       hphI     ndeI    maeIII    aciI
401 ATGTATTTCA GATGCTAAAC ATGAATTCAG CTCCAACTTT CATCAACTTT CCTGCAAAAG GGAAACCCAA ACGGGGTGAT ACATATGAGT TACAGGTGCG
    TACATAAAGT CTACGATTTG TACTTAAGTC GAGGTTGAAA GTAGTTGAAA GGACGTTTTC CCTTTGGGTT TGCCCCACTA TGTATACTCA ATGTCCACGC
127  V  F  Q   M  L  N   M  N  S  A   P  T  F   I  N  F   P  A  K  G   K  P  K   R  G  D   T  Y  E  L   Q  V  R

               ddeI[M.aluI-]
               bspCNI         mspI    sau3AI
              celII/espI      hpaII   mboI/ndeII[dam-]
              blpI/bpu1102I  scrFI[M.hpaII-]                                                        bsmFI
             aluI            nciI    dpnII[dam-]                                                    sau96I
            pvuII            dsaV    dpnI[dam+]                                                     nlaIV
            mspA1I/nspBII    bssKI alwI[dam-]              sspI                           tsp509I   avaII bslI
501 GGGTTTTTCA GCTGAGCAGA TTGCCCGGTG GATCGCCGAC AGAACTGATG TCAATATTAG AGTGATTAGA CCCCCAAATT ATGCTGGTCC CCTTATGTTG
    CCCAAAAAGT CGACTCGTCT AACGGGCCAC CTAGCGGCTG TCTTGACTAC AGTTATAATC TCACTAATCT GGGGGTTTAA TACGACCAGG GGAATACAAC
160  G  F  S   A  E  Q  I   A  R  W   I  A  D   R  T  D  V   N  I  R   V  I  R   P  P  N  Y   A  G  P   L  M  L

                              taqI                                                           aluI
                             sfuI                                                           tseI
                             bstBI                                           fokI           fnu4HI/bsoFI
                             bsiCI                 tsp509I    tru9I          bstF5I         bbvI
                       baeI     mboII mboII        apoI       mseI       bsrI       mwoI hpyCH4V
601 GGATTGCTTT TGGCTGTTAT TGGTGGACTT GTGTATCTTC GAAGAAGTAA TATGGAATTT CTCTTTAATA AAACTGGATG GGCTTTTGCA GCTTTGTGTT
    CCTAACGAAA ACCGACAATA ACCACCTGAA CACATAGAAG CTTCTTCATT ATACCTTAAA GAGAAATTAT TTTGACCTAC CCGAAAACGT CGAAACACAA
193 G  L  L  L   A  V  I   G  G  L   V  Y  L  R   R  S  N   M  E  F   L  F  N  K   T  G  W   A  F  A   A  L  C  F
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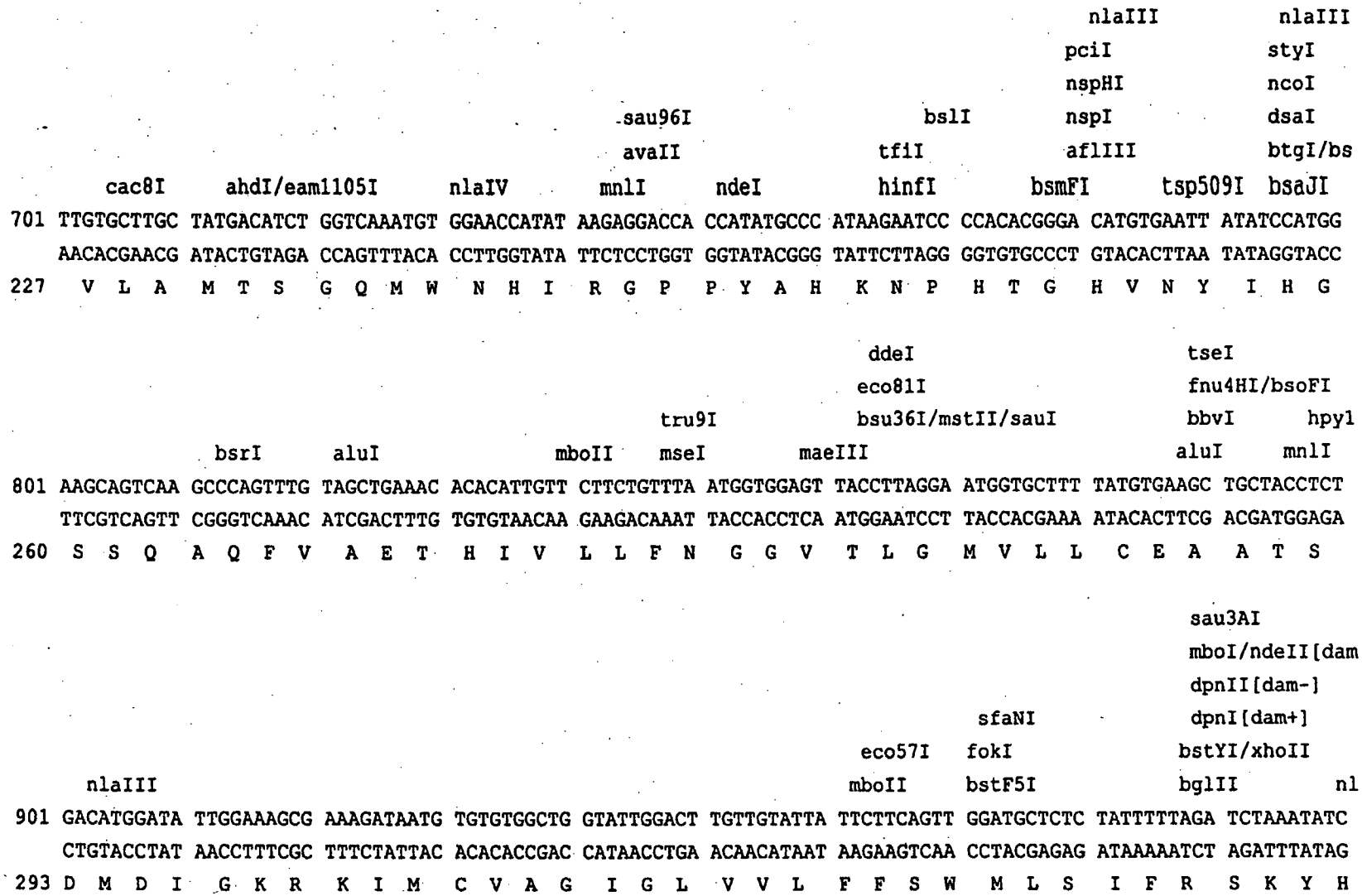
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pciI styI
nspHI ncoI
sau96I bslI nspI dsaI
avaII tfiI aflIII btgI/bs
cac8I ahdI/eam1105I nlaIV mnlI ndeI hinfI bsmFI tsp509I bsaJI
701 TTGTGCTTGC TATGACATCT GGTCAAATGT GGAACCATAT AAGAGGACCA CCATATGCCC ATAAGAATCC CCACACGGGA CATGTGAATT ATATCCATGG
AACACGAACG ATACTGTAGA CCAGTTTACA CCTTGGTATA TTCTCCTGGT GGTATACGGG TATTCTTAGG GGTGTGCCCT GTACACTTAA TATAGGTACC
227 V L A M T S G Q M W N H I R G P P Y A H K N P H T G H V N Y I H G
ddeI tseI
eco81I fnu4HI/bsoFI
tru9I bsu36I/mstII/sauI bbvI hpyI
bsrI aluI mboII mseI maeIII aluI mnlI
801 AAGCAGTCAA GCCCAGTTTG TAGCTGAAAC ACACATTGTT CTTCTGTTTA ATGGTGGAGT TACCTTAGGA ATGGTGCTTT TATGTGAAGC TGCTACCTCT
TTCGTCAGTT CGGGTCAAAC ATCGACTTTG TGTGTAACAA GAAGACAAAT TACCACCTCA ATGGAATCCT TACCACGAAA ATACACTTCG ACGATGGAGA
260 S S Q A Q F V A E T H I V L L F N G G V T L G M V L L C E A A T S
sau3AI
mboI/ndeII[dam
dpnII[dam-]
sfaNI dpnI[dam+]
eco57I fokI bstYI/xhoII
nlaIII mboII bstF5I bglII nl
901 GACATGGATA TTGGAAAGCG AAAGATAATG TGTGTGGCTG GTATTGGACT TGTTGTATTA TTCTTCAGTT GGATGCTCTC TATTTTTAGA TCTAAATATC
CTGTACCTAT AACCTTTCGC TTTCTATTAC ACACACCGAC CATAACCTGA ACAACATAAT AAGAAGTCAA CCTACGAGAG ATAAAAATCT AGATTTATAG
293 D M D I G K R K I M C V A G I G L V V L F F S W M L S I F R S K Y H

```
                                    bsmFI
                                    sau96I                         bsrI
                                    nlaIV                         rsaI
                                    avaII                    bpmI/gsuI[dcm-]
                               tru9I ppuMI                   bsrI  csp6I
                      aluI hpy188I mseI eco0109I/draII        tspRI scaI    tsp509I
1001 ATGGCTACCC ATACAGCTTT CTGATGAGTT AAAAAGGTCC CAGAGATATA TAGACACTGG AGTACTGGAA ATTGAAAAAC GAAAATCGTG TGTGTTTGAA
     TACCGATGGG TATGTCGAAA GACTACTCAA TTTTTCCAGG GTCTCTATAT ATCTGTGACC TCATGACCTT TAACTTTTTG CTTTTAGCAC ACACAAACTT
327   G  Y  P  Y  S  F  L  M  S  O

                                                   tru9I
                                                   mseI  tru9I
      bsmI                                       ahaIII/draI    tru9I                    tru9I
    mboII hpyCH4V                  mnlI          swaI     mseI  mseI     mboII           mseI
1101 AAGAAGAATG CAACTTGTAT ATTTGTATT ACCTCTTTTT TTCAAGTGAT TTAAATAGTT AATCATTTAA CCAAAGAAGA TGTGTAGTGC CTTAACAAGC
     TTCTTCTTAC GTTGAACATA TAAAACATAA TGGAGAAAAA AAGTTCACTA AATTTATCAA TTAGTAAATT GGTTTCTTCT ACACATCACG GAATTGTTCG

                 mnlI
                ddeI                                    bsrI                     tsp509I    tru9I
                bspCNI                mnlI tru9I     mboII tspRI               tru9I    rsaI tsp509I
     mnlI      hpy188I            tsp509I  mseI      earI/ksp632I              mseI     csp6I  mseI
1201 AATCCTCTGT CAAAATCTGA GGTATTTGAA AATAATTATC CTCTTAACCT TCTCTTCCCA GTGAACTTTA TGGAACATTT AATTTAGTAC AATTAAGTAT
     TTAGGAGACA GTTTTAGACT CCATAAACTT TTATTAATAG GAGAATTGGA AGAGAAGGGT CACTTGAAAT ACCTTGTAAA TTAAATCATG TTAATTCATA

                                                             tru9I
                                                             mseI
                                                            hpaI
    psiI  tsp509I                         aluI              hincII/hindII hpy188I              bslI
1301 ATTATAAAAA TTGTAAAACT ACTACTTTGT TTTAGTTAGA ACAAAGCTCA AAACTACTTT AGTTAACTTG GTCATCTGAT TTTATATTGC CTTATCCAAA
     TAATATTTTT AACATTTTGA TGATGAAACA AAATCAATCT TGTTTCGAGT TTTGATGAAA TCAATTGAAC CAGTAGACTA AAATATAACG GAATAGGTTT
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                                     scrFI[dcm-]
                                     pspGI
                                     mvaI
                                     ecoRII[dcm-]
                                     dsaV[dcm-]
                                     bstNI                                                      tsp509I[M.ecoRI-]
                                     bssKI[dcm-]                                               xmnI
                                     apyI[dcm+]                                                ecoRI           mboII
                                    sexAI                                                      asp700      aluI      mslI        fokI
                               hpy188III             ndeI       maeIII                         apoI      ddeI[M.aluI-]           bstF5I
1401 GATGGGGAAA GTAAGTCCTG ACCAGGTGTT CCCACATATG CCTGTTACAG ATAACTACAT TAGGAATTCA TTCTTAGCTT CTTCATCTTT GTGTGGATGT
     CTACCCCTTT CATTCAGGAC TGGTCCACAA GGGTGTATAC GGACAATGTC TATTGATGTA ATCCTTAAGT AAGAATCGAA GAAGTAGAAA CACACCTACA

                                                                                                         taiI
                                                                                                   hgiAI/aspHI
                                                                                                   bsp1286
                                                                          hpy188I                  bsiHKAI    rmaI  ddeI
     bstZ17I                                                            mboII                  hpy188I   maeII/hpyCH4IV
     bst1107I                                                          bpuAI                eco57I    aflIII maeI  bspC
     accI        sfaNI                        tsp509I     nlaIII bbsI                      mboII  bmyI btrI  bfaI mnlI
1501 GTATACTTTA CGCATCTTTC CTTTTGAGTA GAGAAATTAT GTGTGTCATG TGGTCTTCTG AAAATGGAAC ACCATTCTTC AGAGCACACG TCTAGCCCTC
     CATATGAAAT GCGTAGAAAG GAAAACTCAT CTCTTTAATA CACACAGTAC ACCAGAAGAC TTTTACCTTG TGGTAAGAAG TCTCGTGTGC AGATCGGGAG
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                                                                                   tth111I/aspI
                                                                                   pleI
                     bseRI                                                         pflFI
                    mnlI                              bpmI/gsuI[dcm-]              mlyI
                   bseRI                            hinPI       ddeI               hinfI
          bst4CI/hpyCH4III mnlI  hpyCH4V            hhaI/cfoI   bspCNI           bsmAI     bsmAI
1601 AGCAAGACAG TTGTTTCTCC TCCTCCTTGC ATATTTCCTA CTGCGCTCCA GCCTGAGTGA TAGAGTGAGA CTCTGTCTCA AAAAAAAGTA TCTCTAAATA
     TCGTTCTGTC AACAAAGAGG AGGAGGAACG TATAAAGGAT GACGCGAGGT CGGACTCACT ATCTCACTCT GAGACAGAGT TTTTTTTCAT AGAGATTTAT

                                     tru9I                                                                    tsp45I
                                     mseI                              tfiI                                  hphI
             tsp509I               hpaI                   xmnI         hinfI                              tru9I maeIII
         psiI           smlI       hincII/hindII          asp700     hpy188I           ddeI               mseI bstEII
1701 CAGGATTATA ATTTCTGCTT GAGTATGGTG TTAACTACCT TGTATTTAGA AAGATTTCAG ATTCATTCCA TCTCCTTAGT TTTCTTTTAA GGTGACCCAT
     GTCCTAATAT TAAAGACGAA CTCATACCAC AATTGATGGA ACATAAATCT TTCTAAAGTC TAAGTAAGGT AGAGGAATCA AAAGAAAATT CCACTGGGTA

                                                                                                                  dde
                     ddeI[M.aluI-]       maeIII          haeIII/palI                     tsp45I            rsaI
                   aluI             tspRI          nlaIII                    tsp509I     maeIII            csp6I
1801 CTGTGATAAA AATATAGCTT AGTGCTAAAA TCAGTGTAAC TTATACATGG CCTAAAATGT TTCTACAAAT TAGAGTTTGT CACTTATTCC ATTTGTACCT
     GACACTATTT TTATATCGAA TCACGATTTT AGTCACATTG AATATGTACC GGATTTTACA AAGATGTTTA ATCTCAAACA GTGAATAAGG TAAACATGGA
```

```
                                         scrFI[dcm-]
                                         pspGI
                                         mvaI
                                         ecoRII[dcm-]
                                         dsaV[dcm-]
                                         bstNI
                                       haeIII/palI
                                      mscI/balI[dcm-]
                                      eaeI[dcm-]
                                      cfrI
                                    scrFI[dcm-]
                                    pspGI
                                    mvaI bssKI[dcm-]
                                    ecoRII[dcm-]  tsp45I
                                    dsaV[dcm-]    maeIII                                          bssS
                                    bstNI    hinPI                                                hpy18
                                    bssKI[dcm-]  tspRI                                           sau3AI
                              pleI bslI[dcm-] hhaI/cfoI                            styI cac8I    mboI/nd
               ddeI           mlyI bsaJI apyI[dcm+]                ddeI          haeIII/palI     dpnII[d
               bspCNI         hinfI apyI[dcm+]  btsI               bspCNI      mnlI bsaJI        dpnI[da
1901 AAGAGAAAAA TAGGCTCAGT TAGAAAAGGA CTCCCTGGCC AGGCGCAGTG ACTTACGCCT GTAATCTCAG CACTTTGGGA GGCCAAGGCA GGCAGATCAC
     TTCTCTTTTT ATCCGAGTCA ATCTTTTCCT GAGGGACCGG TCCGCGTCAC TGAATGCGGA CATTAGAGTC GTGAAACCCT CCGGTTCCGT CCGTCTAGTG
```

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                                  mscI/ball[dcm-]
                                  eaeI[dcm-]
                                 scrFI[dcm-]
                                 pspGI
                                 mvaI
                                 ecoRII[dcm-]
                                 dsaV[dcm-]
                                 bstNI
                        bsmAI    bssKI[dcm-]
                      taqI   fokI  cfrI    nlaIII         bsmAI
          hpy188III  bsaI bstF5I haeIII/palI          esp3I                    aluI
    mnlI          hpy188III  apyI[dcm+]   hphI       bsmBI                 tsp509I                nlaIV
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     CTCCAGTCCT CAAGCTCTGG TAGGACCGGT TGTACCACTT TGGGGCAGAG ATGATTTTTA TATTTTTAAT CGACCCACAC CACCGTCCTC GGACATTAGG

                                                                                                    scrFI[dcm-]
                                                                                                    pspGI
                                                                                                    mvaI
                                                                                                    ecoRII[dcm-]
                                                                                  tspRI             dsaV[dcm-]
                                                                          sau3AI  btsI              bstNI
                      ddeI         tfiI          hpy188III                mboI/ndeII[dam-]          bssKI[dcm-]
                      bspCNI       hinfI         ddeI                     dpnII[dam-] hpyCH4V       apyI[dcm+]
      aluI       mnlI  mnlI bssSI               bspCNI      mnlI    tspRI  dpnI[dam+] bsgI  bpmI/gsuI[dcm-]
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     GTCGATGTGT CCTCCGACTC CGTGCTCTTA GTGAACTTGA GTCCTCTACC TCCAAAGTCA CTCGGCTCTA GTGCGGTGAC GTGAGGTCGG ACCGTTGTCT
```

```
                                                          fnu4HI/bsoFI
                                                         haeIII/palI
                                                        mcrI
                                                        eagI/xmaIII/eclXI
                                                        eaeI
                                                        cfrI
          pleI                                          bsiEI      rmaI
          mlyI                                        notI         maeI
          hinfI                                       fnu4HI/bsoFI bfaI
        bsmAI                                         aciI aciI  speI
2201 GCGAGACTCC ATCTCAAAAA AAAAAAAAAA AAAAAAAAAA AAAAAAAGGG CGGCCGCCGA CTAGTGAGC
     CGCTCTGAGG TAGAGTTTTT TTTTTTTTTT TTTTTTTTTT TTTTTTCCC GCCGGCGGCT GATCACTCG
```

> length: 2269

accI(GTMKAC):	1501
aciI(CCGC):	39 498 2250 2254
aflIII(ACRYGT):	780 1586
ahaIII(TTTAAA):	1150
ahdI(GACNNNNNGTC):	278 714
aluI(AGCT):	152 300 429 510 690 822 888 1015 1345 1476 1816 2070 2102
alw26I(CAGNNNCTG):	101 316
alwI(GGATCNNNN):	318 530
alwNI(CAGNNNCTG):	101 316
apoI(RAATTY):	3 310 423 655 1464
apyI(CCWGG):	321 332 1422 1934 1939 2023 2189
asp700(GAANNNNTTC):	1464 1749
aspHI(GWGCWC):	1582